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FULL ESTIMATED COST
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FILE 'BIOSIS' ENTERED AT 14:59:48 ON 16 DEC 2002
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COPYRIGHT (C) 2002 AMERICAN CHEMICAL SOCIETY (ACS)
=> s caryopsis(w)specific(w)promoter
L1
             2 CARYOPSIS(W) SPECIFIC(W) PROMOTER
=> d l1 1-2 ibi ab
'IBI' IS NOT A VALID FORMAT FOR FILE 'CAPLUS'
The following are valid formats:
ABS -----GI and AB
ALL ----- BIB, AB, IND, RE
APPS ----- AI, PRAI
BIB ----- AN, plus Bibliographic Data and PI table (default)
CAN ----- List of CA abstract numbers without answer numbers
CBIB ----- AN, plus Compressed Bibliographic Data
DALL ----- ALL, delimited (end of each field identified)
DMAX ----- MAX, delimited for post-processing
FAM ----- AN, PI and PRAI in table, plus Patent Family data
FBIB ----- AN, BIB, plus Patent FAM
IND ----- Indexing data
IPC ----- International Patent Classifications
MAX ----- ALL, plus Patent FAM, RE
PATS ----- PI, SO
SAM ----- CC, SX, TI, ST, IT
SCAN ------ CC, SX, TI, ST, IT (random display, no answer numbers;
             SCAN must be entered on the same line as the DISPLAY,
             e.g., D SCAN or DISPLAY SCAN)
STD ----- BIB, IPC, and NCL
IABS ----- ABS, indented with text labels
IALL ----- ALL, indented with text labels
IBIB ----- BIB, indented with text labels
IMAX ----- MAX, indented with text labels
ISTD ----- STD, indented with text labels
OBIB ----- AN, plus Bibliographic Data (original)
OIBIB ----- OBIB, indented with text labels
SBIB ----- BIB, no citations
SIBIB ----- IBIB, no citations
HIT ----- Fields containing hit terms
HITIND ----- IC, ICA, ICI, NCL, CC and index field (ST and IT)
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containing hit terms HITRN ----- HIT RN and its text modification HITSTR ----- HIT RN, its text modification, its CA index name, and its structure diagram HITSEQ ----- HIT RN, its text modification, its CA index name, its structure diagram, plus NTE and SEO fields FHITSTR ---- First HIT RN, its text modification, its CA index name, and its structure diagram FHITSEQ ---- First HIT RN, its text modification, its CA index name, its structure diagram, plus NTE and SEQ fields KWIC ----- Hit term plus 20 words on either side OCC ----- Number of occurrence of hit term and field in which it occurs To display a particular field or fields, enter the display field codes. For a list of the display field codes, enter HELP DFIELDS at an arrow prompt (=>). Examples of formats include: TI; TI, AU; BIB, ST; TI, IND; TI, SO. You may specify the format fields in any order and the information will be displayed in the same order as the format specification. All of the formats (except for SAM, SCAN, HIT, HITIND, HITRN, HITSTR, FHITSTR, HITSEQ, FHITSEQ, KWIC, and OCC) may be used with DISPLAY ACC to view a specified Accession Number. ENTER DISPLAY FORMAT (BIB): ibib ANSWER 1 OF 2 CAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 2002:31660 CAPLUS DOCUMENT NUMBER: 136:80964 TITLE: Caryopsis-specific promoter of wheat for use in tissue-specific expression of foreign genes in cereal INVENTOR (S): Sprunck, Stephanie; Kluth, Antje; Becker, Dirk; Luetticke, Stephanie; Loerz, Horst PATENT ASSIGNEE(S): Aventis Cropscience Gmbh, Germany SOURCE: PCT Int. Appl., 64 pp. CODEN: PIXXD2 DOCUMENT TYPE: Patent LANGUAGE: German FAMILY ACC. NUM. COUNT: PATENT INFORMATION: PATENT NO. KIND DATE APPLICATION NO. DATE ---------A1 20020110 WO 2001-EP7593 WO 2002002786 20010703 W: AE, AG, AL, AM, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CN, CO, CR, CU, CZ, DM, DZ, EC, EE, GD, GE, HR, HU, ID, IL, IN, IS, JP, KG, KP, KR, KZ, LC, LK, LR, LT, LV, MA, MD, MG, MK, MN, MX, NO, NZ, PL, RO, RU, SG, SI, SK, TJ, TM, TT, UA, US, UZ, VN, YU, ZA, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG DE 10032379 DE 2000-10032379 20000706 **A1** 20020117 AU 2001089621 **A5** 20020114 AU 2001-89621 20010703 PRIORITY APPLN. INFO.: DE 2000-10032379 A 20000706 WO 2001-EP7593 W 20010703 REFERENCE COUNT: 7 THERE ARE 7 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L1 ANSWER 2 OF 2 CAPLUS COPYRIGHT 2002 ACS
ACCESSION NUMBER: 2002:31659 CAPLUS
DOCUMENT NUMBER: 136:80963
TITLE: A caryopsis-specific

promoter of wheat for use in the

tissue-specific expression of foreign genes in cereal

INVENTOR(S): Sprunck, Stefanie; Kluth, Antje; Becker, Dirk;

Luetticke, Stephanie; Loerz, Horst Aventis Cropscience Gmbh, Germany

SOURCE: PCT Int. Appl., 57 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent German

LANGUAGE:
FAMILY ACC. NUM. COUNT:

2

7

PATENT INFORMATION:

PATENT ASSIGNEE(S):

PATENT NO. KIND DATE APPLICATION NO. DATE ----------WO 2002002785 A1 20020110 WO 2001-EP7592 20010703 W: AE, AG, AL, AM, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CN, CO, CR, CU, CZ, DM, DZ, EC, EE, GD, GE, HR, HU, ID, IL, IN, IS, JP, KG, KP, KR, KZ, LC, LK, LR, LT, LV, MA, MD, MG, MK, MN, MX, NO, NZ, PL, RO, RU, SG, SI, SK, TJ, TM, TT, UA, US, UZ, VN, YU, ZA, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG DE 10032379 DE 2000-10032379 20000706 A1 20020117 DE 10041861 A120020314 DE 2000-10041861 20000826 AU 2001077523 A5 20020114 AU 2001-77523 20010703 PRIORITY APPLN. INFO.: DE 2000-10032379 A 20000706 DE 2000-10041861 A 20000826

WO 2001-EP7592 W 20010703
THERE ARE 7 CITED REFERENCES AVAILABLE FOR THIS

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AND TECHNOLOGY CORPORATION, AND FACHINFORMATIONSZENTRUM KARLSRUHE

FILE CONTAINS CURRENT INFORMATION.
LAST RELOADED: Dec 13, 2002 (20021213/UP).

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COST IN U.S. DOLLARS

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FULL ESTIMATED COST

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=> s caryopsis(w)promoter

PRAI DE 2000-10032379 A

DE 2000-10041861 A

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=> d l2 1-2
L2
      ANSWER 1 OF 2 CAPLUS COPYRIGHT 2002 ACS
AN
      2002:31660 CAPLUS
DN
      136:80964
      Caryopsis-specific promoter of wheat for use in tissue-specific expression
ΤI
      of foreign genes in cereal
      Sprunck, Stephanie; Kluth, Antje; Becker, Dirk; Luetticke, Stephanie;
IN
      Loerz, Horst
PA
      Aventis Cropscience Gmbh, Germany
SO
      PCT Int. Appl., 64 pp.
      CODEN: PIXXD2
DT
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LA
      German
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TI
      A caryopsis-specific promoter of wheat for use in the tissue-specific
      expression of foreign genes in cereal
      Sprunck, Stefanie; Kluth, Antje; Becker, Dirk; Luetticke, Stephanie;
IN
      Loerz, Horst
PA
      Aventis Cropscience Gmbh, Germany
SO
      PCT Int. Appl., 57 pp.
      CODEN: PIXXD2
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FAN.CNT 2
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L4
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     136:80964
ΤI
     Caryopsis-specific promoter of wheat for use in
     tissue-specific expression of foreign genes in cereal
     Sprunck, Stephanie; Kluth, Antje; Becker, Dirk; Luetticke, Stephanie;
IN
     Loerz, Horst
PA
     Aventis Cropscience Gmbh, Germany
SO
     PCT Int. Appl., 64 pp.
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AN
    2002:31659 CAPLUS
DN
    136:80963
    A caryopsis-specific promoter of wheat for use in the
TI
    tissue-specific expression of foreign genes in cereal
IN
    Sprunck, Stefanie; Kluth, Antje; Becker, Dirk; Luetticke, Stephanie;
    Loerz, Horst
PA
    Aventis Cropscience Gmbh, Germany
SO
    PCT Int. Appl., 57 pp.
    CODEN: PIXXD2
DT
    Patent
    German
LA
FAN.CNT 2
    PATENT NO.
                   KIND DATE
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WO 2001-EP7592

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AN
     1957:53959 CAPLUS
DN
     51:53959
OREF 51:9993f-q
     The influence of interring straw and of nitrogen fertilization on the
     quality of caryopses of the succeeding wheat crop
ΑU
     Cavazza, Luigi
CS
     Univ. Bari, Italy
SO
     Ann. sper. agrar. (Rome) (1957), 11, 25-34
DT
     Journal
LA
     English
=> s caryopsis and specific and promoter
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PROCESSING COMPLETED FOR L5
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L6
     ANSWER 1 OF 5 CAPLUS COPYRIGHT 2002 ACS
     A starch-associated R1 protein of Solanum tuberosum and its use in
     altering starch properties of transgenic wheat and other plants
L6
     ANSWER 2 OF 5 CAPLUS COPYRIGHT 2002 ACS
TI
     Caryopsis-specific promoter of wheat for use
     in tissue-specific expression of foreign genes in cereal
L6
     ANSWER 3 OF 5 CAPLUS COPYRIGHT 2002 ACS
ΤI
     A caryopsis-specific promoter of wheat for
     use in the tissue-specific expression of foreign genes in cereal
     ANSWER 4 OF 5 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.DUPLICATE 1
L6
     Functional characterization of seed coat-specific members of the
     barley germin gene family.
L6
    ANSWER 5 OF 5 CAPLUS COPYRIGHT 2002 ACS
TI
     Gene expression using fusion with promoter of crop plant lipid
     transfer protein gene
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=> d 16 1 4 5 ibib ab

L6 ANSWER 1 OF 5 CAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 2002:332351 CAPLUS DOCUMENT NUMBER: 136:322182

TITLE:

A starch-associated R1 protein of Solanum tuberosum and its use in altering starch properties of

transgenic wheat and other plants

INVENTOR (S):

Schewe, Gabi; Knies, Petra; Amati, Simone Franceska;

Loerz, Horst; Becker, Dirk; Uwer, Ursula;

Landschuetze, Volker; Pilling, Jens; Frohberg, Klaus

Aventis Cropscience Gmbh, Germany

PATENT ASSIGNEE(S): SOURCE:

PCT Int. Appl., 81 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

FAMILY ACC. NUM. COUNT:

English

PATENT INFORMATION:

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PATENT NO.
                   KIND DATE
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    WO 2002034923 A2 20020502
                                      WO 2001-EP12179 20011022
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            LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PH, PL,
            PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG,
            UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
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PRIORITY APPLN. INFO.:
                                     DE 2000-10052492 A 20001023
                                     DE 2000-10064805 A 20001222
                                     WO 2001-EP12179 W 20011022
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The present invention relates to monocotyledon plant cells and plants AΒ which are genetically modified to encode a starch-assocd. R1 protein. Plant cells and plants of this type synthesize a modified starch, which has an increased phosphate content and/or a modified phosphorylation pattern and/or an increased final viscosity in an RVA profile and/or a reduced peak temp. in differential scanning calorimetry anal. and/or an increased gel strength in the texture anal. compared with starch from corresponding non-genetically modified monocotyledon plants. specifically, the transgenic plant cell synthesized a starch with a phosphate at the C6 position of a glucose monomer and has a phosphate content of at least 0.1 nmol phosphate/mg of starch. However, the amylose component of starch has a reduced total phosphate content. Furthermore, the modified starch has a 50% increase in viscosity and a peak temp. that is reduced by at least 1.5.degree.C. Therefore, the present invention also relates to the starch which is synthesized from the plant cells and plants according to the invention, and to methods of producing said starch. The present invention further relates to wheat flours which contain said modified starches, and to food products and bakery products which contain said wheat flours and/or starch.

ANSWER 4 OF 5 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.DUPLICATE 1

ACCESSION NUMBER: DOCUMENT NUMBER:

2001:49506 BIOSIS PREV200100049506

TITLE:

Functional characterization of seed coat-specific

members of the barley germin gene family.

AUTHOR (S):

Wu, Shiping; Druka, Arnis; Horvath, Henriette; Kleinhofs, Andris; Kannangara, C. Gamini; von Wettstein, Diter (1)

CORPORATE SOURCE:

(1) Departments of Crop and Soil Sciences and Genetics and Cell Biology, Washington State University, Pullman, WA,

99164: diter@wsu.edu USA

SOURCE:

Plant Physiology and Biochemistry (Paris), (September,

2000) Vol. 38, No. 9, pp. 685-698. print.

ISSN: 0981-9428.

DOCUMENT TYPE:

LANGUAGE:

Article English

SUMMARY LANGUAGE: English

AB The present project aimed to isolate testa-, pericarp- and epicarp-

specific gene promoters for the developing caryopsis of barley (Hordeum vulgare L.). These might be applied in transgenic plants to express antifungal agents or modify metabolic pathways. A testa-specific 379-nucleotide fragment was cloned by differential amplification and used to screen a bacterial artificial chromosome (BAC) library of 6.3 haploid genome equivalents. Fifty-three clones containing genes encoding for proteins of the germin family were found. Characterization of the clones identified a minimum of six seed coat- and eight leaf-specific germin genes. Four seed coat- and one leaf-specific genes were sequenced. The deduced primary structure of the proteins revealed a remarkable conservation of the manganese(II) binding His and Glu residues and beta-barrel secondary structure of oxalate oxidase - also in barley, wheat, rice and Arabidopsis germins, for which an enzymatic activity has not yet been identified. The oxalate oxidase and germins of barley and other species are synthesized with a conserved pre-sequence of 23 or 24 amino acids for targeting into the cell wall. beta-Glucuronidase expression with the barley germin F gene promoter occurs specifically in the testa and epicarp of the developing barley caryopsis, while expression with the B gene promoter is restricted to the testa. Oxalate oxidase activity is prominent in the epicarp and the root tips of the developing embryo. A family tree based on primary structure homologies of germins distinguishes three groups: oxalate oxidases, leaf-specific germins and seed coat-specific germins.

L6 ANSWER 5 OF 5 CAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1995:907813 CAPLUS

DOCUMENT NUMBER:

123:308197

TITLE:

Gene expression using fusion with promoter of crop plant lipid transfer protein gene

INVENTOR(S):

Olsen, Odd-Arne; Kalla, Roger; Linnestad, Casper

PATENT ASSIGNEE(S):

Norway

SOURCE:

PCT Int. Appl., 64 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent English

LANGUAGE:

n Eng.

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

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PATENT NO.
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PRIORITY APPLN. INFO.:
                                    GB 1994-3512
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AB An expression system for at least the aleurone cells of a developing caryopsis or for at least the scutellar epithelial tissue or vascular tissue of a germinating seedling or developing grain or plant (e.g. in the root, leaves and stem) is described. The expression system comprises a gene promoter fused to a GOI (gene of interest). In a preferred embodiment the expression system comprises the GOI fused to a modified Ltp 1 gene promoter.

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- TI Matrix attachment regions.
- AU Jordan, Mark Carlyle (1); Rampitsch, Christof; Cloutier, Marie Sylvie Jacqueline
- CS (1) Winnipeg Canada

 ASSIGNEE: Her Majesty the Queen in right of Canada, as represented by the
 Department of Agriculture and Agri-Food Canada, Lethbridge, Canada
- PI US 6177612 January 23, 2001
- SO Official Gazette of the United States Patent and Trademark Office Patents,

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(Jan. 23, 2001) Vol. 1242, No. 4, pp. No Pagination. e-file.
      ISSN: 0098-1133.
 DT
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      Endosperm-specific activity of a storage protein gene
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      Lamacchia, C.; Shewry, P.R.; Di Fonzo, N.; Forsyth, J.L.; Harris, N.;
      Lazzeri, P.A.; Napier, J.A.; Halford, N.G.; Barcelo, P.
      DNAL (450 J8224)
      Journal of experimental botany, Feb 2001. Vol. 52, No. 355. p. 243-250
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      Publisher: Oxford : Oxford University Press.
      CODEN: JEBOA6; ISSN: 0022-0957
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ΑU
     Cloutier, S.; Rampitsch, C.; Penner, G.A.; Lukow, O.M.
ΑV
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     Publisher: London ; New York : Academic Press, c1983-
     CODEN: JCSCDA; ISSN: 0733-5210
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     Endosperm-specific GFP expression driven by barley
     D-hordein promoter and its inheritance in transgenic barley and
     wheat plants.
     Cho, M.-J. (1); Kim, H.-K. (1); Choi, H. W. (1); Buchanan, B. B. (1);
ΑU
     Lemaux, P. G. (1)
     (1) Department of Plant and Microbial Biology, University of California,
     Berkeley, CA, 94720 USA
     In Vitro Cellular & Developmental Biology Animal, (March, 2000) Vol. 36,
     No. 3 Part 2, pp. 63.A. print.
     Meeting Info.: Meeting of the Society for In Vitro Biology World Congress
     on In Vitro Biology San Diego, California, USA June 10-15, 2000
     ISSN: 1071-2690.
DT
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LA
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L10
     2000:60182 AGRICOLA
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     Overexpression of thioredoxin h leads to enhanced activity of starch
ΤI
     debranching enzyme (pullulanase) in barley grain.
ΑU
     Cho, M.J.; Wong, J.H.; Marx, C.; Jiang, W.; Lemaux, P.G.; Buchanan, B.B.
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      Publisher: Washington, D.C.: National Academy of Sciences,
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      1999:485703 CAPLUS
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      131:267754
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AU
     Li, Z.; Rahman, S.; Kosar-Hashemi, B.; Mouille, G.; Appels, R.; Morell, M.
CS
     CSIRO Plant Industry, Canberra, 2601, Australia
SO
     Theoretical and Applied Genetics (1999), 98(8), 1208-1216
     CODEN: THAGA6; ISSN: 0040-5752
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     Springer-Verlag
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     Journal
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     An endosperm-specific DOF protein from barley, highly
     conserved in wheat, binds to and activates transcription from the
     prolamin-box of a native B-hordein promoter in barley
     endosperm.
ΑU
     Mena, M.; Vicente-Carbajosa, J.; Schmidt, R.J.; Carbonero, P.
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     ETS Ingenieros Agronomos, Madrid, Spain.
ΑV
     DNAL (QK710.P68)
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     1. p. 53-62
     Publisher: Oxford : Blackwell Sciences Ltd.
     ISSN: 0960-7412
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     97:45195 AGRICOLA
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     The wheat transcriptional activator SPA: a seed-specific bZIP protein that
TI
     recognizes the GCN4-like motif in the bifactorial endosperm box
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AU
     Albani, D.; Hammond-Kosack, M.C.U.; Smith, C.; Conlan, S.; Colot, V.;
     Holdsworth, M.; Bevan, M.W.
CS
     John Innes Centre, Norwich, UK.
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     Publisher: [Rockville, MD : American Society of Plant Physiologists,
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ANSWER 9 OF 12 AGRICOLA

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      promoter in plant and yeast cells.
      Holdsworth, M.J.; Munoz-Blanco, J.; Hammond-Kosack, M.; Colot, V.; Schuch,
 ΑU
      W.; Bevan, M.W.
 CS
      John Innes Centre, Norwich, UK.
      DNAL (QK710.P62)
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      Publisher: Dordrecht : Kluwer Academic Publishers.
      CODEN: PMBIDB; ISSN: 0167-4412
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      93:41937 AGRICOLA
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      In vivo footprinting of a low molecular weight glutenin gene (LMWG-1D1) in
     wheat endosperm.
     Hammond-Kosack, M.C.U.; Holdsworth, M.J.; Bevan, M.W.
ΑU
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     John Innes Centre for Plant Sciences Research, Norwich, UK
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     Structural and functional analysis of promoter from gliadin, an
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     Triticum aestivum L.
ΑU
     Aryan, A.P.; An, G.; Okita, T.W.
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     Washington State University, Pullman, WA
ΑV
     DNAL (442.8 Z34)
     M G G: Molecular and general genetics, Jan 1991. Vol. 225, No. 1. p.
     65-71 ill
     Publisher: Berlin, W. Ger. : Springer International.
     CODEN: MGGEAE; ISSN: 0026-8925
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L10 ANSWER 12 OF 12 AGRICOLA
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     92:32091 AGRICOLA
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     Identification of an enhancer element for the endosperm-
     specific expression of high molecular weight glutenin.
     Thomas, M.S.; Flavell, R.B.
CS
     University of Nottingham Medical School, Nottingham, United Kingdom
ΑV
     DNAL (QK725.P532)
     The Plant cell, Dec 1990. Vol. 2, No. 12. p. 1171-1180
     Publisher: Rockville, Md. : American Society of Plant Physiologists.
     ISSN: 1040-4651
NTE Includes references.
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- DTArticle
- FS U.S. Imprints not USDA, Experiment or Extension
- English
- => d l10 1-12 ab
- ANSWER 1 OF 12 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.DUPLICATE
- Matrix attachment regions isolated from the 5' flanking region of endosperm-specific storage protein genes of monocotyledonous plants are provided. An exemplified matrix attachment region is derived from the 5' flanking region of the Bx7 gluten gene of Triticum aestivum. Recombinant nucleic acid molecules and plant vectors containing such recombinant nucleic acid molecules include DNA constructs having a promoter, a coding sequence, and a poly(A) addition site, the DNA constructs operably linked to at least one of the matrix attachment regions. Gene expression in transgenic plants, preferably monocotyledonous cereal crop species, is improved by transforming plants with such recombinant nucleic acid molecules.
- ANSWER 2 OF 12 AGRICOLA

DUPLICATE 2

- The characterization of the promoter of a wheat (Triticum aestivum) cv. Cheyenne high molecular weight glutenin subunit (HMW subunit) gene, Glu-1D-1 is reported. The nucleotide sequence of the promoter from position -1191 to -650 with respect to the transcription start site was determined, to add to that already determined. Analysis of this region of the promoter revealed the presence of an additional copy of part of the primary enhancer sequence and sequences related to regulatory elements present in other wheat seed protein genes. A chimaeric gene was constructed comprising the 5' flanking region of the Glu-1D-1 gene from position -1191 to +58, the coding region of the UidA (Gus) gene, and the nopaline synthase (Nos) gene terminator. This chimaeric gene was introduced into wheat (Triticum durum cv. Ofanto) by particle bombardment of inflorescence explants. Two independent transgenic lines were produced, and both showed expression of the Gus gene specifically in the endosperm during mid-development (first detected 10-12 d after anthesis). Histochemical analysis of homozygous T2 seed confirmed this pattern of expression, and showed that expression was initiated first in the central lobes of the starchy endosperm, and then spread throughout the endosperm tissue, while no expression was detected in the aleurone layer. Native HMW subunit protein was detectable by Western analysis 12-14 d after anthesis, consistent with concurrent onset of activity of the native and introduced HMW subunit gene promoters.
- L10 ANSWER 3 OF 12 AGRICOLA
- AB Endosperm-specific low-molecular-weight (LMW) glutenins are an important component of the polymeric gluten and, as such, play a key role in end-use functionality. Reports of N-terminal amino acid sequences of LMW glutenin fractions revealed that they have either a methionine or a serine residue at the first position of the mature peptide. These subunits were therefore called LMW-m and LMW-s type glutenins. A gene that is predicted to encode a LMW glutenin subunit having an isoleucine amino acid residue at position one of the mature protein was amplified and cloned from extra strong bread wheat cultivar Glenlea (pGH3.1). The predicted N-terminal sequence of this gene is truncated as compared to the m-type and s-type. The gene still codes for the expected number of eight cysteine residues which are all located in the C-terminal region. We propose to call it LMW-i based on the same nomenclature. Analysis of 277 doubled haploid lines derived from a single cross showed perfect co-segregation of the cloned PCR fragment with a rare LMW glutenin called LMW-50. The gene was subcloned in an expression vector and the protein was expressed in E. coli. Western blot analysis using a

prolamin-specific monoclonal antibody confirmed the co-migration of the cloned protein with LMW-50 from Glenlea.

- L10 ANSWER 4 OF 12 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.
- L10 ANSWER 5 OF 12 AGRICOLA
- Biochemically active wheat thioredoxin h has been overexpressed in the endosperm of transgenic barley grain. Two DNA constructs containing the wheat thioredoxin h gene (wtrxh) were used for transformation; each contained wtrxh fused to an endospermspecific B(1)-hordein promoter either with or without a signal peptide sequence for targeting to the protein body. Twenty-two stable, independently transformed regenerable lines were obtained by selecting with the herbicide bialaphos to test for the presence of the bar herbicide resistance gene on a cotransformed plasmid; all were positive for this gene. The presence of wtrxh was confirmed in 20 lines by PCR analysis, and the identity and level of expression of wheat thioredoxin h was assessed by immunoblots. Although levels varied among the different transgenic events, wheat thioredoxin h was consistently highly expressed (up to 30-fold) in the transgenic grain. Transgenic lines transformed with the B(1)-hordein promoter with a signal peptide sequence produced a higher level of wheat thioredoxin h on average than those without a signal sequence. The overexpression of thioredoxin h in the endosperm of germinated grain effected up to a 4-fold increase in the activity of the starch debranching enzyme, pullulanase (limit dextrinase), the enzyme that specifically cleaves alpha-1,6 linkages in starch. These results raise the question of how thioredoxin h enhances the activity of pullulanase because it was found that the inhibitor had become inactive before the enzyme showed appreciable activity.
- L10 ANSWER 6 OF 12 CAPLUS COPYRIGHT 2002 ACS
- A cDNA clone, and a corresponding genomic DNA clone, contg. full-length sequences encoding wheat starch synthase I, were isolated from a cDNA library of hexaploid wheat (Triticum aestivum) and a genomic DNA library of Triticum tauschii, resp. The entire sequence of the starch synthase-I cDNA (wSSI-cDNA) is 2591 bp, and it encodes a polypeptide of 647 amino-acid residues that shows 81% and 61% identity to the amino-acid sequences of SSI-type starch synthases from rice and potato, resp. In addn., the putative N-terminal amino-acid sequence of the encoded protein is identical to that detd. for the N-terminal region of the 75-kDa starch synthase present in the starch granule of hexaploid Two prominent starch synthase activities were demonstrated to be present in the sol. fraction of wheat endosperm by activity staining of the non-denaturing PAGE gels. The most anodal band (wheat SSI) shows the highest staining intensity and results from the activity of a 75-kDa protein. The wheat SSI mRNA is expressed in the endosperm during the early to mid stages of wheat grain development but was not detected by Northern blotting in other tissues from the wheat plant. The gene encoding the wheat SSI (SsI-D1) consists of 15 exons and 14 introns, similar to the structure of the rice starch synthase-I gene. While the exons of wheat and rice are virtually identical in length, the wheat SsI-D1 gene has longer sequences in introns 1, 2, 4 and 10, and shorter sequences in introns 6, 11 and 14, than the corresponding rice gene.
- L10 ANSWER 7 OF 12 AGRICOLA
- AB A cDNA encoding a DNA-binding protein of the DOF class of transcription factors was isolated from a barley endosperm library. The deduced amino acid sequence for the corresponding protein is 94% identical through the DOF domain to the prolamin-box (P-box) binding factor PBF from maize. The gene encoding the barley PBF (BPBF) maps to chromosome 7H, and its expression is restricted to the endosperm where it precedes that of the hordein genes. The BPBF expressed in bacteria as a GST-fusion binds a P-box 5'-TGTAAAG-3' containing oligonucleotide derived from the

promoter region of an Hor2 gene. Binding was prevented when the P-box motif was mutated to 5-'TGTAgAc-3'. A P-box binding activity, present in barley and wheat endosperm nuclei, interacted similarly to BPBF with this synthetic oligonucleotide, and the binding was abolished by 1,10-phenanthroline. Transient expression experiments in developing barley endosperms demonstrate that BPBF transactivates transcription from the P-box element of a native Hor2 promoter and that direct binding of BPBF to its target site is essential for transactivation since mutations in the DOF DNA-binding domain or in the P-box motif of this promoter abolished both binding and transactivation. Evidence was also obtained for the presence in wheat of a Pbf homologue having similar DNA-binding properties to that of BPBF. These results strongly implicate this endospermspecific DOF protein from barley as an important activator of hordein gene expression and suggest the evolutionary conservation of the Pbf gene function among small grain cereals.

L10 ANSWER 8 OF 12 AGRICOLA

The conserved bifactorial endosperm box found in the promoter of wheat storage protein genes comprises two different cis elements that are thought to be involved in regulating endosperm-specific gene expression. Endosperm nuclear extracts contain binding activities. One is called ESBF-I, which binds to the endosperm motif (EM), and the other is called ESBF-II, which binds to the GCN4-like motif (GLM). Here, we present a functional analysis of the endosperm box of a low-molecular-weight glutenin gene found on the 1D1 chromosome of hexaploid wheat (LMWG-1D1) in transgenic tobacco plants. Our analysis demonstrates the necessity of the EM and GLM for endospermspecific gene expression and suggests the presence in tobacco of functional counterparts of wheat ESBF-I and ESBF-II. Furthermore, we describe the isolation and characterization of cDNA clones encoding SPA, a seed-specific basic leucine zipper protein from wheat that can activate transcription from the GLMs of the -326-bp LMWG-1D1 promoter in both maize and tobacco leaf protoplasts. This activation is also partially dependent on the presence of functional EMs, suggesting interactions between SPA with ESBF-I-like activities.

L10 ANSWER 9 OF 12 AGRICOLA

The **promoter** of the wheat low-molecular-weight glutenin AB (LMWG1D1) gene contains a cis element called the GCN4 like motif (GLM) which has low homology to one class of binding site for the maize endosperm-specific b-ZIP transcription factor Opaque-2 (O2). Previous work has shown that the GLM element interacts with the nuclear factor ESBFII during wheat endosperm development at the time of maximum transcription of the LMWG1D1 gene. In this paper we demonstrate that O2 binds to the GLM element and can activate high levels of transcription from the wheat GLM in transient assays in plant protoplasts and in yeast cells. Lower levels of O2 activation through the GLM element in yeast containing a defective GCN4 gene showed that GCN4 was necessary for high levels of O2 transcriptional activation, indicating that O2 may need to heterodimerise with GCN4 to activate transcription in yeast. These observations provide evidence that the GLM represents a new type of O2 DNA-binding site, and support a postulate that an O2 homologue may activate endosperm-specific expression of wheat storage protein genes.

L10 ANSWER 10 OF 12 AGRICOLA

The quality of the wheat grain is determined by the quantity and composition of storage proteins (prolamins) which are synthesized exclusively in endosperm tissue. We are investigating the mechanisms underlying the regulation of expression of a prolamin gene, the low molecular weight glutenin gene LMWG-1D1. The LMWG-1D1 promoter contains the endosperm box, a sequence motif highly conserved in

the promoter region of a large number of storage protein genes, which is thought to confer endosperm-specific expression of prolamin genes. Here we show by in vivo DMS footprinting of wheat endosperm tissue that the endosperm box becomes occupied by putative trans-acting factors during grain ripening. During early stages of development the endosperm motif within the 5' half of the endosperm box becomes occupied first, followed by binding of a second activity to a GCN4/jun-like motif in the 3' half just prior to the stage of maximum gene expression. Occupancy of the endosperm box is highly tissue-specific: no protection was observed in husk and leaf tissues. Several binding activities were identified in vitro from nuclear protein extracts of wheat endosperm which bind specifically to the endosperm and GCN4/jun motifs identified by in vivo footprinting.

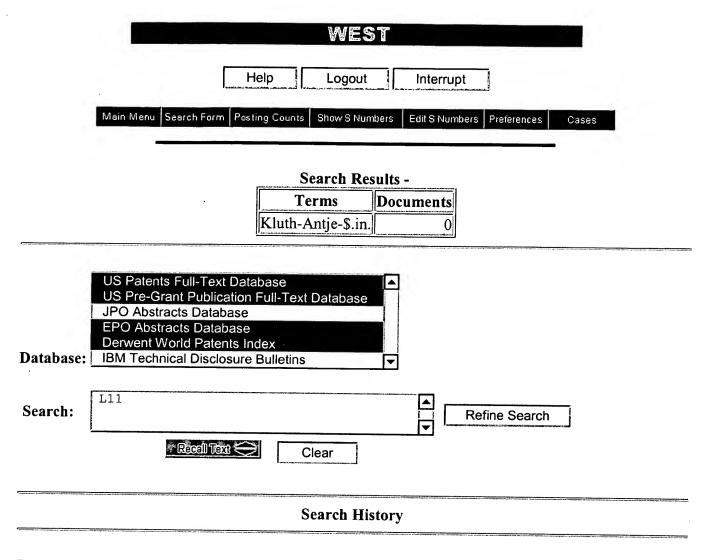
L10 ANSWER 11 OF 12 AGRICOLA

DUPLICATE 3 To identify cis-regulatory elements of the gliadin gene, a study of the AB gliadin gene promoter was conducted by transient expression analysis of plasmid DNAs which were introduced into plant protoplasts by electroporation. The **promoter** region (-592 bp to +18 bp from the translational start) of this developmentally regulated gene, when fused upstream to the chloramphenicol acetyl transferase (CAT) reporter cassette was unable to direct significant CAT expression in wheat or tobacco suspension cells. Because this monocot gene promoter appeared to be under stringent tissue-specific control, a hybrid promoter approach using a nopaline synthase (nos) promoter was employed. A series of 3' deletions of the gliadin promoter were placed upstream of either a nonfunctional -101 nos or a nearly wild-type -155 nos promoter fused in turn to a CAT reporter gene cassette. Transient expression analysis of these plasmid DNAs in tobacco cells showed that the gliadin fragment could either restore the activity of the non-functional nos promoter (series I) or enhance the activity of the functional nos promoter (series II). The degree of restoration of the promoter function conferred by gliadin fragments of the first series was proportional to the enhancing effect of the same fragments in the second series of constructs. The transcriptional activity of the gliadin (-592 bp to -77 bp) -nos hybrid promoter was reduced by 26% upon 3' deletion of sequences in the region -141 bp to -77 bp, which contains both the TATA and CCAAT boxes. A marked decline in the promoter function of these hybrid constructs, however, was observed when an additional upstream region was removed, suggesting the presence of regulatory sequences in the -218 bp to -141 bp region of the gliadin promoter. Deletion of the -300 bp element, which is similar to the SV40 core enhancer, did not affect hybrid promoter function, although additional upstream activating sequences (-592 bp to -448 bp) were also observed.

L10 ANSWER 12 OF 12 AGRICOLA

Genes encoding high molecular weight (HMW) glutenin, a wheat seed storage protein, are expressed only in the developing endosperm. It was previously demonstrated that sequences essential for endospermspecific transcription reside within 436 base pairs upstream of the initiation codon for HMW glutenin translation. We have further analyzed this region by testing the ability of a series of truncated HMW glutenin promoter fragments to enhance transcription from an adjacent heterologous promoter. The activity of these hybrid promoters was determined by measuring the expression of a linked beta-glucuronidase (GUS) reporter gene in transgenic tobacco plants. An HMW glutenin promoter fragment spanning nucleotides -375 to -45 relative to the transcription start site was found to stimulate GUS expression in tobacco seeds when inserted in either orientation upstream of the heterologous promoter. Furthermore, this fragment could also potentiate transcription when located 3' to the GUS reporter gene. Stimulation of GUS gene expression in transgenic tobacco seeds did not

occur until 9 days to 12 days after anthesis, coincident with the onset of storage protein synthesis in the developing tobacco and wheat seed, and was confined to the **endosperm** tissue. By testing progressively shorter **promoter** fragments, the enhancer element responsible for this pattern of expression was localized to a 40-base pair region some 170 base pairs upstream of the start site for HMW glutenin transcription.



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END OF SEARCH HISTORY